

002220" 6F202560

1/11

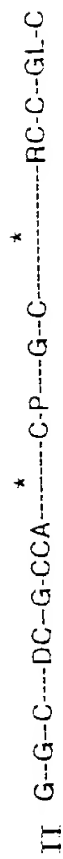
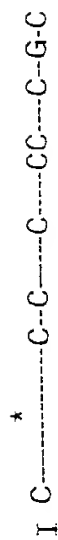


Fig. 1

1	6C6GTG6C66CC	✓	phdtk-3	2.1	✓
1	✓	pcdtk-3	2.2	✓
1	✓	pndtk-2	2.3	✓
1	✓	phdtk-2	2.4	✓
1	6ACCCACGCTCCGTGCTTGTGCTTCG	✓	pndtk-1	2.5	✓
1	TGGCCCCCGCACGCAAAATTC	✓	phdtk-1	2.6	✓
1	6ACA6TCG6AGCCGCGCTGCAAGCATCAAGGACTTATCTTGGAGGACTTGTGAATTC	✓	pRNdtk-1	2.7	✓
13	6CTCTAGAAATAGTGGATCCCCCGGCTGCAAGAAATTCGGCACGAGCGGCTGCGGCGCGCA6	✓	phdtk-3		
1	✓	pcdtk-3		
1	✓	pndtk-2		
1	✓	phdtk-2		
34	GA6ATGATGGTTGTGTGTGCAcCGGCA6CTGTCCGGTTCCTGGCCGTGTTACAAATGATG	✓	pndtk-1		
23	GGCAGcGAGGGTCTGgGcgcTcagagGATGCTCTGACCTTGAAAGGGTCCCTATCTGGAGAGCG	✓	phdtk-1		
61	CATCCTGCCATTGTGGTTACTGAGTCTGCTGGTTGGACAGAGGAAAGGGCAGCAACATGTTCC	✓	pRNdtk-1		
73	AGCGGAGATGCAAGCGGCTTGGGGCCCAACCCTGCTGTGCTGCTGGCGGCGGCGGCTCCG	✓	phdtk-3		
4	CGAGCGGCAAGCGGCTGAGAGCGCGCGGCGATGCGGCGGCGGAGAGGAGCGGCGGCGGCG	✓	pcdtk-3		
1	✓	pndtk-2		
1	✓	phdtk-2		
94	GCTCTCTGCAAGCCTCCCTCTGCTAGGAGCCAGTGGCCACCTTGAACTCA6TTC...TCATCA	✓	pndtk-1		
83	AGGAGGTACAAACGTGCTGAAATGTTGTGCGGTTTCAGGGAGGCAATTTGGTTAACCCCT...GCCATT	✓	phdtk-1		
121	C6GTGCCCTCTTATTGTCTTTTGGGGYTTTATCTTGGATG6G6GCACTTGGCTTTGTGATGA	✓	pRNdtk-1		

Fig. 2

[illegible]

Fig. 2 (Forts.)

004220" 6T202560

312	CTGCTA	AAAGCA	ATCATCA	GAAAGT	GAAACC	TGGCAAA	CTTACC	CTCCAG	CTAT	CACAA	TGAGAA	TGAGAA
244	GGGCAAA	AAAAAC	TGTCA	GAAAGT	AAACTT	TGAAAA	AACTTACC	CTCCAC	CTTACC	CTTACC	CTTACC	CTTACC
106	GGCACAG	TC	CC
67	GGCACAG	TC	CC
329	GCTCCAG	CC	CC	AA	6CCG	CGG	CGG	CGG	CGG	CGG	CGG	CGG
314	GGCTAG	TC	CC
361	GGCACAG	TC	CC
372	CCAA	CACAG	AC	GAAG	GT	GGAA	ATAA	TA	TA	TA	TA	TA
304	CCAA	CACAG	AA	CA	GA	AT	GGTA	ATAA	TA	TA	TA	TA
148	GGAA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA
109	GGAA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA
389	GA	AA	GC	CA	GG	AA	GC	CA	GG	AA	GC	CA
371	GGAA	GC	CG	CA	AA	GC	CA	GG	AA	GC	CG	CA
406	GGAA	AC	GC	CA	GA	AA	GC	CA	GG	AA	GC	CA
432	T
364	T	AC	GA	TA	AC	GA	AA	CT	GG	AT	CA	CA
208	AT	GG	AA	AT	CT	GG	CA	TC	CC	CA	CT	GG
169	AT	GG	CA	AT	CT	GG	CA	TC	CC	CA	CT	GG
449	AT	GG	AA	AT	CT	GG	CA	TC	CC	CA	CT	GG
431	AT	GG	AA	AT	CT	GG	CA	TC	CC	CA	CT	GG
466	AC	GG	AA	AT	CT	GG	CA	TC	CC	CA	CT	GG

phdtk-3
pcdtk-3
pmdtk-2
phdtk-2
pmdtk-1
phdtk-1
pRNdtk-1

phdtk-3
pcdtk-3
pmdtk-2
phdtk-2
pmdtk-1
phdtk-1
pRNdtk-1

phdtk-3
pcdtk-3
pmdtk-2
phdtk-2
pmdtk-1
phdtk-1
pRNdtk-1

Fig. 2 (Forts.)

	phd1dk-3	pcd1dk-3	pmc1dk-2	phd1dk-2	pmc1dk-1	phd1dk-1	pmc1dk-1
433	GTGGAGAAACAAAGAA	TCATGAGTG	ATCATTTGA	TGAAGAC	TGTGAAC	AAACAAG	AAAGT
424	ATGGCAACCCGGCATAGAG	ATGGTCACTA	TTCTCAACCA	TGACCTGGG	ATGGC	ATGGC	ATGGC
265	ATGGTACTCGGCACAGAG	ATGGTCACTA	TTCTCAACCA	TGACCTGGG	ATGGC	ATGGC	ATGGC
226	ATGGTACTCGGCACAGAG	ATGGTCACTA	TTCTCAACCA	TGACCTGGG	ATGGC	ATGGC	ATGGC
500	AGGAAAGCATCATTTGAAG	ATGGTCACTA	TTCTCAACCA	TGACCTGGG	ATGGC	ATGGC	ATGGC
479	AGGAAAGCATCATTTGAAG	ATGGTCACTA	TTCTCAACCA	TGACCTGGG	ATGGC	ATGGC	ATGGC
526	AGGAAAGCATCATTTGAAG	ATGGTCACTA	TTCTCAACCA	TGACCTGGG	ATGGC	ATGGC	ATGGC

[illegible]

	phdtkc-3	pcdtkc-3	pmdtkc-2	phdtkc-2	pmdtkc-1	phdtkc-1	p2Ndktkc-1
433	GC	TC	CA	CG	AT	GT	TT
544	GC	TC	CA	CG	AT	GT	TT
385	GC	TC	CA	CG	AT	GT	TT
346	GC	TC	CA	CG	AT	GT	TT
620	GC	TC	CA	CG	AT	GT	TT
593	GC	TC	CA	CG	AT	GT	TT
643	GC	TC	CA	CG	AT	GT	TT

Fig. 2 (Forts.)

433
 783 CCTGATGGAATACTAGAGCGCTGCCCATGTGCAAGTGGCTTGAATCTGGCCAACTTCAAGAGC
 619 AAACACTGGAAGAGTCAATCAGTACAGACCTGAGAAATTTGGTGAATTTAAAGCAATTAAGGC
 580 CACCAATTGAGGAACATCAATCAATGCAAGACCTGAGAAATTTGGTGAATTTAAAGCAATTAAGGC
 856 ... ACCGACAGTC...TAAATATGATGGACTCTTTTATCTAAATAATGCTACGAAATATC
 829
 883 CGAGGCTACAGAG...CCTGAAGGACCTTCTCTAAATTAAGCTAAATTAAGACTTTGGTATC
 433
 843 AGCCACACATGTAATCTGTTGTTGAACTGCTCCCTCCAAATGAAACCAAGCAAGAAACCAAGAA
 678 ATGATGGAAGAACCTGGATTTGGAATGCGGAAGAAATGAGGGAATGTTGGAATGTTGGAGCAAG
 640 ATGATGGAAGAAATGAAGTTCAAGATGCAAGAAAGAAAT...GGCTAAAAATGAAGAAACGTTGAATGAAG
 910 CTTTATGATTTGTCAGCTCAATCCCAAGGATGTAAGGAATCTTCAAGTGGTGAATTAAGCAAT
 829
 941 CTGCAATGTTATTTTCTCAATTTTACATGAAGTGGCTCTGGTCTTCCCTGAAACCCGGAAGTGGCTG
 433
 903 GAAGATCCCTTGAACATGGATGAGATGCCCATTTATCAATTTAAATACCCAGAGAGATATCTCT
 738 AAGAGGCGAGGACTGAAATCAAGTAGAGTCCGACAAACCAAGAAAGTACTACCAAGTGGCTTCCG
 697 AATATAGATGATCACAAGAAAGAAAGAAAGAAAGATGCGGCGCAAGCTTATTTCCCTTTA
 970 TCCGACAAATACTTTCCAAAGGCTCTGGAAGTGTGAAGGACTTTGGTTCTTGAATGGAAGCTCC
 829
 1001 CGCAACTTGTCTTTTCTTGTGAGGAACTTCCCTAAATTAAGCTAAATTAAGCTAAATTAAGCT

phdtkk-3
 pcdtkk-3
 pmcdtkk-2
 phdtkk-2
 pmcdtkk-1
 phdtkk-1
 pRNdtkk-1

phdtkk-3
 pcdtkk-3
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 phdtkk-1
 pRNdtkk-1

phdtkk-3
 pcdtkk-3
 pmcdtkk-2
 phdtkk-2
 pmcdtkk-1
 phdtkk-1
 pRNdtkk-1

Fig. 2 (Forts.)

[illegible]

Fig. 2 (Forts.)

Fig. 2 (Forts.)

[illegible]

433	phdckc-3
1323	C T T G T A C A G C A G A A T A A C G T A T C A G T A C T C A T T A A A A A C A C A C G G A G C A	phdckc-3
882	phdckc-2
769	phdckc-2
1227	phdckc-1
829	phdckc-1
1298	phdckc-1

10/11

453	.	phdckc-3
1303	T	pcdckc-3
882	.	pmckc-2
769	.	phdckc-2
1227	.	pmckc-1
829	.	phdckc-1
1290	.	piRNckckc-1

Fig. 2 (Forts.)

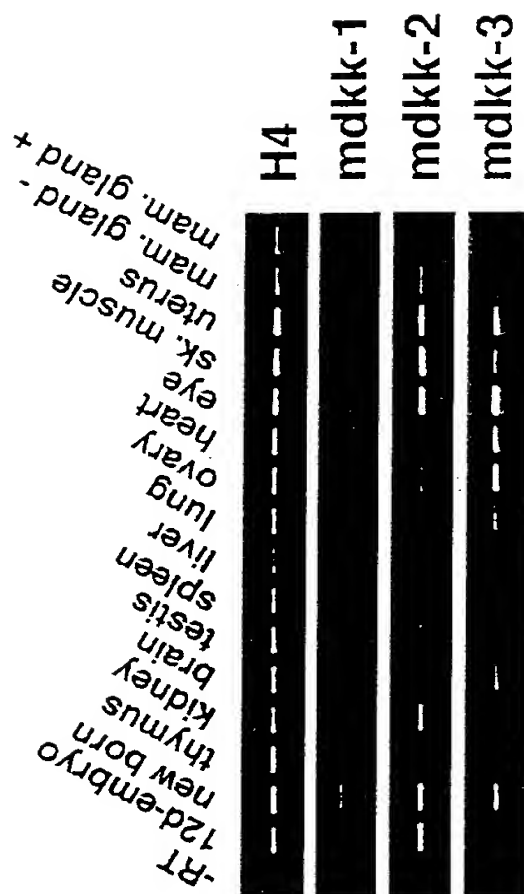


Fig. 3